AMENDMENTS TO THE CLAIMS:

Claims 16, 18-47 and 49-56 are canceled without prejudice or disclaimer. Claims 57-74 are added. The following is the status of the claims of the above-captioned application, as amended.

Claims 1-56 (Canceled).

N76D+S103A+V104L

Claim 57 (New). A variant of a parent subtilase comprising a substitution at position 9 with Arg and a substitution at position 62 with Asp, wherein the variant has protease activity and each position corresponds to a position of the amino acid sequence of SEQ ID NO: 1.

Claim 58 (New). The variant of claim 57, further comprising a substitution at position 68.

Claim 59 (New). The variant of claim 58, wherein the substitution at position 68 is V68A.

Claim 60 (New). The variant of claim 57, further comprising one or more of the modifications K27R, "36D, S56P, V68A, N76D, S87N, G97N, S99SE, S101G, S103A, V104A, V104I, V104N, V104Y, S106A, H120D, H120N, N123S, G159D, Y167A, R170L, R170S, A194P, N204D, V205I, Q206E, L217D, N218S, N218D, M222S, M222A, T224S, A232V, K235L, Q236H, Q245R, N248D, N252K, T274A, S3T+V4I+S99D+S101R+S103A+V104I+G160S+A194P+V199M+V205I+L217D, S3T+V4I+S99D+S101R+S103A+V104I+G160S+V199M+V205I+L217D, S3T+V4I+S99D+S101R+S103A+V104I+G160S+V199M+V205I+L217D, S3T+V4I+S99D+S101R+S103A+V104I+G160S+V205I. K27R+V104Y+N123S+T274A,

N76D+V104A

S87N+S101G+V104N

S99D+S101R+S103A+V104I+G160S, and S101G+V104N.

Claim 61 (New). The variant of claim 57, further comprising the following substitutions: S101G+S103A+V104I+G159D+A232V+Q236H+Q245R+N248D+N252K.

Claim 62 (New). The variant of claim 57, comprising a set of substitutions selected from the group consisting of:

S9R+A13V+A15T+I35V+N62D+Q245F;
S9R+A15T+G20*+L21F+P52T+N62D+Q245R;
,
S9R+A15T+G20*+L21F+N62D+Q245N;
S9R+A15T+G20*+L21F+N62D+Q245R; and
S9R+A15T+G20*+L21F+N62D+Q245R+S259G.

Claim 63 (New). The variant of claim 57, comprising a set of substitutions selected from the group consisting of:

S9R+A15T+T22A+N62D;
S9R+A15T+T22TG+N62D+V139L+Q245G;
S9R+A15T+T22TG+N62D+V139L+Q245S;
S9R+A15T+T22TG+N62D+V139L+Q245V;
S9R+A15T+T22TL+N62D+I107V+V139L+Q245W; and
S9R+A15T+T22TL+N62D+Q245W.

Claim 64 (New). The variant of claim 57, comprising a set of substitutions selected from the group consisting of:

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$9R+A15T+N62D+H120N+P131T;
$9R+A15T+N62D+Q245R+N252M;
$9R+A15T+N62D+Q245W+N252S; and
$9R+A15T+N62D+Q245W+N252V.
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Claim 65 (New). The variant of claim 57, wherein the parent subtilase belongs to the subgroup I-S1.

Claim 66 (New). The variant of claim 57, wherein the parent subtilase belongs to the subgroup I-S2. Claim 67 (New). The variant of claim 57, wherein the parent subtilase is subtilisin 309.

Claim 68 (New). A cleaning or detergent composition, comprising a variant of claim 57 and a surfactant.

Claim 69 (New). A composition of claim 68, which additionally comprises one or more of an amylase, cellulase, cutinase, esterase, beta-galactosidase, glycoamylase, hemicellulase, lactase, ligninase, lipase, polygalacturonase, and protease.

Claim 70 (New). An isolated DNA sequence encoding a variant of claim 57.

Claim 71 (New). An expression vector comprising the isolated DNA sequence of claim 70.

Claim 72 (New). A microbial host cell transformed with the expression vector of claim 71.

Claim 73 (New). A microbial host cell of claim 72, which is a Bacillus.

Claim 74 (New). A method for producing a variant, comprising

- (a) culturing a host of claim 72 under conditions conducive to the expression and secretion of the variant, and
 - (b) recovering the protease variant.